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SEQUENCE LISTING

<110> Frohberg, Claus Koetting, Oliver

> Ritte, Gerhard Steup, Martin

- <120> Plants with increased activity of multiple starch phosphorylating enzymes
- <130> 65084.000022
- <140> Unassigned <141> 2006-09-01
- <150> EP 04090089.6
- <151> 2004-03-05
- <150> US 60/550,021
- <151> 2004-03-05
- <150> EP 04090284.3
- <151> 2004-07-21
- <150> EP 04090121.7
- <151> 2004-03-29
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Glu Asp Ala Ser Ala Ser Val Thr Phe Ala Pro Glu Lys Val Gln Asp Page 17 195 200 205

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450

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Gly Lys Phe Asn Leu Glu Gly Asn Val Glu Leu Gln Ile Thr Val Gly 100 105 110

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His Asn Asn Thr Ser Pro Asp Asp Val Ile Ile Cys Gln Ala Leu Ile Page 31

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Leu Ser Asp Asp Ile Asn Gln Ala Val Ala Glu Lys Leu Gln Ile Page 33 1205 1210 1215

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Val Leu Thr Pro Asp Met Pro Asp Val Leu Ser His Val Ser Val	Arg 3069
yal Leu Thr Pro Asp Met Pro Asp Val Leu Ser His Val Ser Val 995 1000 1005 gca aga aat gga aag atc tgc ttt gcc aca tgt ttt gat tct ggl Ala Arg Asn Gly Lys Ile Cys Phe Ala Thr Cys Phe Asp Ser Gly	Arg 3069
yal Leu Thr Pro Asp Met Pro Asp Val Leu Ser His Val Ser Val 995 1000 1005 gca aga aat gga aag atc tgc ttt gcc aca tgt ttt gat tct ggt Ala Arg Asn Gly Lys Ile Cys Phe Ala Thr Cys Phe Asp Ser Gly 1010 1015 1020 atc tta tct gac ctc caa gga aaa gat gga aaa ctg ttg agc cta Ile Leu Ser Asp Leu Gln Gly Lys Asp Gly Lys Leu Leu Ser Leu	Arg 3069 y 3114 u 3159
Val Leu Thr Pro Asp Met Pro Asp Val Leu Ser His Val Ser Val 995 1000 1005 gca aga aat gga aag atc tgc ttt gcc aca tgt ttt gat tct ggt Ala Arg Asn Gly Lys Ile Cys Phe Ala Thr Cys Phe Asp Ser Gly 1010 1015 1020 atc tta tct gac ctc caa gga aaa gat gga aaa ctg ttg agc cta Ile Leu Ser Asp Leu Gln Gly Lys Asp Gly Lys Leu Leu Ser Leu 1025 1030 1035 caa cca acc tct gca gat gta gtc tat aaa gag gta aac gat agt Gln Pro Thr Ser Ala Asp Val Val Tyr Lys Glu Val Asn Asp Ser	Arg 3069 y 3114 u 3159 r 3204
Val Leu Thr Pro Asp Met Pro Asp Val Leu Ser His Val Ser Val 995 gca aga aat gga aag atc tgc ttt gcc aca tgt ttt gat tct ggt Ala Arg Asn Gly Lys Ile Cys Phe Ala Thr Cys Phe Asp Ser Gly 1010 atc tta tct gac ctc caa gga aaa gat gga aaa ctg ttg agc cta Ile Leu Ser Asp Leu Gln Gly Lys Asp Gly Lys Leu Leu Ser Leu 1025 caa cca acc tct gca gat gta gtc tat aaa gag gta aac gat agc Gln Pro Thr Ser Ala Asp Val Val Tyr Lys Glu Val Asn Asp Ser 1040 gag ctt tcg agt cca agt tca gac aac ctg gaa gat gcc cct cca Glu Leu Ser Ser Pro Ser Ser Asp Asn Leu Glu Asp Ala Pro Pro	Arg t 3069 a 3114 t 3159 r 3204 a 3249
Val Leu Thr Pro Asp Met Pro Asp Val Leu Ser His Val Ser Val 995 gca aga aat gga aag atc tgc ttt gcc aca tgt ttt gat tct ggt Ala Arg Asn Gly Lys Ile Cys Phe Ala Thr Cys Phe Asp Ser Gly 1010 atc tta tct gac ctc caa gga aaa gat gga aaa ctg ttg agc cta Ile Leu Ser Asp Leu Gln Gly Lys Asp Gly Lys Leu Leu Ser Leu 1025 caa cca acc tct gca gat gta gtc tat aaa gag gta Asn Asp Ser Gly Pro Thr Ser Ala Asp Val Val Tyr Lys Glu Val Asn Asp Ser Gly Leu Ser Ser Pro Ser Ser Asp Asn Leu Glu Asp Ala Pro Pro 1055 agt att tct ttg gtc aag aaa cag ttt gcg ggt aga tat gct atc Ser Ile Ser Leu Val Lys Lys Gln Phe Ala Gly Arg Tyr Ala Ile	Arg 3069 y 3014 u 3159 r 3204 a 3249 e 3294

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gtg Val	atc Ile 1385	aga Arg	gac Asp	ggg Gly	aag Lys	ctc Leu 1390	tat Tyr	gtc Val	gtc Val	cag Gln	aca Thr 1395	cga Arg	cca Pro	caa Gln	4194
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Tyr Gly Lys Gly Leu Lys Gln Ala Gly Arg Ser Leu Val Thr Glu Thr 50 55 60

Gly Gly Arg Pro Leu Ser Phe Val Pro Arg Ala Val Leu Ala Met Asp 65 70 75 80

Pro Gln Ala Ala Glu Lys Phe Ser Leu Asp Gly Asn Ile Asp Leu Leu $85 \hspace{1cm} 90 \hspace{1cm} 95$

Val Glu Val Thr Ser Thr Thr Val Arg Glu Val Asn Ile Gln Ile Ala 100 105 110

Tyr Thr Ser Asp Thr Leu Phe Leu His Trp Gly Ala Ile Leu Asp Asn 115 120 125

Lys Glu Asn Trp Val Leu Pro Ser Arg Ser Pro Asp Arg Thr Gln Asn 130 135 140

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Ser	His	Leu	Lys	Leu 165	Glu	Ile	Asp	Asp	Pro 170	Ala	Ile	His	Ala	Ile 175	Glu
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Gln	Asn	Phe 195	His	Ile	Asn	Leu	Pro 200	Thr	Glu	Arg	Asn	Va1 205	Lys	Gln	Asn
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Lys 305	Ile	Gln	Arg	Lys	Gly 310	Arg	Asp	Leu	Thr	Lys 315	Leu	Ile	Tyr	Lys	ніs 320
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485 490 495 Leu Leu Asp Lys Ile Ala Asp Leu Glu Ser Glu Ala Gln Lys Ser Phe 500 505 510 Met His Arg Phe Asn Ile Ala Ala Asp Leu Val Asp Glu Ala Lys Ser 515 520 525 Ala Gly Gln Leu Gly Phe Ala Gly Ile Leu Val Trp Met Arg Phe Met 530 540 Ala Thr Arg Gln Leu Val Trp Asn Lys Asn Tyr Asn Val Lys Pro Arg 545 550 555 560 Glu Ile Ser Lys Ala Gln Asp Arg Leu Thr Asp Leu Leu Gln Asp Val 565 570 575 Tyr Ala Ser Tyr Pro Glu Tyr Arg Glu Leu Leu Arg Met Ile Met Ser 580 585 590 Thr Val Gly Arg Gly Glu Gly Asp Val Gly Gln Arg Ile Arg Asp 595 605 Glu Ile Leu Val Ile Gln Arg Lys Asn Asp Cys Lys Gly Gly Ile Met 610 620 Glu Glu Trp His Gln Lys Leu His Asn Asn Thr Ser Pro Asp Asp Val 625 635 Val Ile Cys Gln Ala Leu Met Asp Tyr Ile Lys Ser Asp Phe Asp Leu 650

Ser Val Tyr Trp Lys Thr Leu Asn Asp Asn Gly Ile Thr Lys Glu Arg Leu Leu Ser Tyr Asp Arg Ala Ile His Ser Glu Pro Asn Phe Arg Gly Glu Gln Lys Asp Gly Leu Leu Arg Asp Leu Gly His Tyr Met Arg Thr Leu Lys Ala Val His Ser Gly Ala Asp Leu Glu Ser Ala Ile Gln Asn 705 710 720 Cys Met Gly Tyr Gln Asp Asp Gly Glu Gly Phe Met Val Gly Val Gln 725 730 735 Ile Asn Pro Val Ser Gly Leu Pro Ser Gly Tyr Pro Asp Leu Leu Arg 740 745 750 Phe Val Leu Glu His Val Glu Glu Lys Asn Val Glu Pro Leu Leu Glu 755 760 765 Gly Leu Leu Glu Ala Arg Gln Glu Leu Arg Pro Leu Leu Leu Lys Ser 770 780 His Asp Arg Leu Lys Asp Leu Leu Phe Leu Asp Leu Ala Leu Asp Ser 785 790 795 800 Thr Val Arg Thr Ala Ile Glu Arg Gly Tyr Glu Gln Leu Asn Asp Ala 805 810 815 Gly Pro Glu Lys Ile Met Tyr Phe Ile Ser Leu Val Leu Glu Asn Leu 820 825 830 Ala Leu Ser Ser Asp Asp Asp Glu Asp Leu Ile Tyr Cys Leu Lys Gly 835 840 Trp Gln Phe Ala Leu Asp Met Cys Lys Ser Lys Lys Asp His Trp Ala 850 860 Leu Tyr Ala Lys Ser Val Leu Asp Arg Ser Arg Leu Ala Leu Ala Ser 865 870 880 Lys Ala Glu Arg Tyr Leu Glu Ile Leu Gln Pro Ser Ala Glu Tyr Leu 885 890 895 Gly Ser Cys Leu Gly Val Asp Gln Ser Ala Val Ser Ile Phe Thr Glu 905

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											Met	. Ser	Asr	n Ser	
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Leu G	ly Asn	Asn	Leu	Leu 10	Tyr	GIn	Gly	Phe	Leu 15	Thr	Ser	Thr	۷a۱	Leu 20	
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Glu H	is Lys	Ser	Arg 25	IIe	Ser	Pro	Pro	30	vaı	GIY	GIY	ASN	Ser 35	Leu	
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	gct Ala															2660
	aag Lys															2708

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Ser Pro Val	Glu Ala Va 1015 cag aat ga	l Gly Tyr Val 102 a atc tac gag	Val Val Val Asp 20 1 aag ccc acg atc 1 Lys Pro Thr Ile	Ğlü Leü 1025 tta gta 3227
Ser Pro Val ctt tca gtt Leu Ser Val gca aaa tct	cag aat ga Gln Asn Gl 1030 qtt aaa gg	l Gly Tyr Val 102 a atc tac gag u Ile Tyr Glu 103 a gag gag gaa	Val Val Val Asp 20 g aag ccc acg atc Lys Pro Thr Ile 35 att cct gat ggt I Ile Pro Asp Gly	Ğlü Leü 1025 tta gta 3227 Leu Val 1040 gct gtt 3272
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<212> PRT

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Gly Asn Ser Leu Phe Gln Gln Gln Val Ile Ser Lys Ser Pro Leu Ser 35 40 45 Thr Glu Phe Arg Gly Asn Arg Leu Lys Val Gln Lys Lys Lys Ile Pro 50 60 Met Glu Lys Lys Arg Ala Phe Ser Ser Ser Pro His Ala Val Leu Thr 65 70 75 80 Thr Asp Thr Ser Ser Glu Leu Ala Glu Lys Phe Ser Leu Gly Gly Asn 85 90 95 Ile Glu Leu Gln Val Asp Val Arg Pro Pro Thr Ser Gly Asp Val Ser 100 105 110Phe Val Asp Phe Gln Val Thr Asn Gly Ser Asp Lys Leu Phe Leu His 115 120 125 Trp Gly Ala Val Lys Phe Gly Lys Glu Thr Trp Ser Leu Pro Asn Asp 130 135 140 Arg Pro Asp Gly Thr Lys Val Tyr Lys Asn Lys Ala Leu Arg Thr Pro 145 150 155 160 Phe Val Lys Ser Gly Ser Asn Ser Ile Leu Arg Leu Glu Ile Arg Asp 165 170 175 Thr Ala Ile Glu Ala Ile Glu Phe Leu Ile Tyr Asp Glu Ala His Asp 180 185 190 Lys Trp Ile Lys Asn Asn Gly Gly Asn Phe Arg Val Lys Leu Ser Arg 195 200 205 Lys Glu Ile Arg Gly Pro Asp Val Ser Val Pro Glu Glu Leu Val Gln 210 215 220 Ile Gln Ser Tyr Leu Arg Trp Glu Arg Lys Gly Lys Gln Asn Tyr Pro 225 230 235 240 Pro Glu Lys Glu Lys Glu Glu Tyr Glu Ala Arg Thr Val Leu Gln 245 250 255 Glu Glu Ile Ala Arg Gly Ala Ser Ile Gln Asp Ile Arg Ala Arg Leu 260 265 270 Thr Lys Thr Asn Asp Lys Ser Gln Ser Lys Glu Glu Pro Leu His Val 275 280 285

Thr Lys Ser Asp Ile Pro Asp Asp Leu Ala Gln Ala Gln Ala Tyr Ile 290 295 300 Arg Trp Glu Lys Ala Gly Lys Pro Asn Tyr Pro Pro Glu Lys Gln Ile 305 310 315 320 Glu Glu Leu Glu Glu Ala Arg Arg Glu Leu Gln Leu Glu Leu Glu Lys Gly Ile Thr Leu Asp Glu Leu Arg Lys Thr Ile Thr Lys Gly Glu Ile 340 345 350 Lys Thr Lys Val Glu Lys His Leu Lys Arg Ser Ser Phe Ala Val Glu 355 360 365 Arg Ile Gln Arg Lys Lys Arg Asp Phe Gly His Leu Ile Asn Lys Tyr 370 375 380 Thr Ser Ser Pro Ala Val Gln Val Gln Lys Val Leu Glu Glu Pro Pro 385 390 395 400 Ala Leu Ser Lys Ile Lys Leu Tyr Ala Lys Glu Lys Glu Glu Gln Ile 405 410 415 Asp Asp Pro Ile Leu Asn Lys Lys Ile Phe Lys Val Asp Asp Gly Glu 420 425 430 Leu Leu Val Leu Val Ala Lys Ser Ser Gly Lys Thr Lys Val His Leu
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440
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Cys	Phe 1085	Asp	Pro	Asn	Ile	Leu 1090	Ala	Asp	Leu	Gln	Ala 1095	Lys	Glu	Gly
Arg	Ile 1100	Leu	Leu	Leu	Lys	Pro 1105	Thr	Pro	Ser	Asp	11e 1110	Ile	Tyr	Ser
Glu	val 1115	Asn	Glu	Ile	Glu	Leu 1120	Gln	Ser	Ser	Ser	Asn 1125	Leu	val	Glu
Ala	Glu 1130	Thr	Ser	Ala	Thr	Leu 1135	Arg	Leu	val	Lys	Lys 1140	Gln	Phe	Gly
Gly	Cys 1145	Tyr	Ala	Ile	Ser	Ala 1150	Asp	Glu	Phe	Thr	Ser 1155	Glu	Met	val
Gly	Ala 1160	Lys	Ser	Arg	Asn	Ile 1165	Ala	Tyr	Leu	Lys	Gly 1170	Lys	٧a٦	Pro
Ser	Ser 1175	val	Gly	Ile	Pro	Thr 1180	Ser	val	Ala	Leu	Pro 1185	Phe	Gly	Val
Phe	Glu 1190	Lys	val	Leu	Ser	Asp 1195	Asp	Ile	Asn	Gln	Gly 1200	٧a٦	Ala	Lys
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Ala	Tyr 1280	Phe	Ser	Thr	Arg	Lys 1285	val	Lys	Leu	Asp	ніs 1290	Asp	Tyr	Leu

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Phe Val Ile His Thr Thr Asn Pro Ser Ser Gly Asp Asp Ser Glu 1310 1315 1320

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190	195	200

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960 965 970

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